

## SEQUENCE LISTING

<110> Serum Biomedical Institute  
<120> METHOD OF PRODUCING RECOMBINANT DNA MOLECULES  
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<150> US 60/480581  
<151> 2003-06-20  
<150> US 60/493586  
<151> 2003-08-07  
<160> 31  
<170> PatentIn version 3.1  
  
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<213> Homo sapiens  
  
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<223> exon #1 of human  $\beta$ -FSH (NM\_000510)

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Met Lys Thr Leu Gln Phe Phe Phe Leu Phe Cys Cys Trp Lys Ala Ile  
1 5 10 15

tgc tgc aat agc tgt gag ctg acc aac atc acc att gca ata gag aaa 96  
Cys Cys Asn Ser Cys Glu Leu Thr Asn Ile Thr Ile Ala Ile Glu Lys  
20 25 30

gaa gaa tgt cgt ttc tgc ata agc atc aac acc act tgg tgt gct ggc 144  
Glu Glu Cys Arg Phe Cys Ile Ser Ile Asn Thr Thr Trp Cys Ala Gly  
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tac tgc tac acc agg  
Tyr Cys Tyr Thr Arg 159  
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acc ttc aag gaa ctg gta tat gaa aca gtg aga gtg ccc ggc tgt gct 96  
 Thr Phe Lys Glu Leu Val Tyr Glu Thr Val Arg Val Pro Gly Cys Ala  
 20 25 30

cac cat gca gat tcc ttg tat aca tac cca gtg gcc acc cag tgt cac 144  
 His His Ala Asp Ser Leu Tyr Thr Tyr Pro Val Ala Thr Gln Cys His  
 35 40 45

tgt ggc aag tgt gac agc gac agc act gat tgt act gtg cga ggc ctg 192  
 Cys Gly Lys Cys Asp Ser Asp Ser Thr Asp Cys Thr Val Arg Gly Leu  
 50 55 60

ggg ccc agc tac tgc tcc ttt ggt gaa atg aaa gaa taa 231  
 Gly Pro Ser Tyr Cys Ser Phe Gly Glu Met Lys Glu  
 65 70 75

<210> 5  
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 5 10 15

Cys Cys Asn Ser Cys Glu Leu Thr Asn Ile Thr Ile Ala Ile Glu Lys  
 20 25 30

Glu Glu Cys Arg Phe Cys Ile Ser Ile Asn Thr Thr Trp Cys Ala Gly  
 35 40 45

Tyr Cys Tyr Thr Arg Asp Leu Val Tyr Lys Asp Pro Ala Arg Pro Lys  
 50 55 60

Ile Gln Lys Thr Cys Thr Phe Lys Glu Leu Val Tyr Glu Thr Val Arg  
 65 70 75 80

Val Pro Gly Cys Ala His His Ala Asp Ser Leu Tyr Thr Tyr Pro Val  
 85 90 95

Ala Thr Gln Cys His Cys Gly Lys Cys Asp Ser Asp Ser Thr Asp Cys  
 100 105 110

Thr Val Arg Gly Leu Gly Pro Ser Tyr Cys Ser Phe Gly Glu Met Lys  
 115 120 125

Glu

<210> 6  
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 <308> GenBank / NM\_000510  
 <309> 2002-11-05  
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 atcaacacca cttgggtgtgc tggctactgc tacaccaggg atctgggtgttaaaggaccca 120  
 gccaggccca aaatccagaa aacatgtacc ttcaaggaac tggtatatga aacagtgaga 180  
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<210> 7  
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<210> 8  
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<210> 9  
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<223> primer PRX2 n.t. position 429-407 in SEQ ID NO: 1, cDNA sequence		
for human $\beta$ -FSH		
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atgcagctat ctttctggtc acattgtcgg tgtttctgca tgttctccat tccgctccctg		180

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 ctttcctcct tattcctacag tacaatcagc agtctagttc ttttcatttg gaatgaatac 660  
 agcattaagc ttgttccact gcaaataaaag cttttaaat catc 704

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 <223> PCR primer HCG-SENT

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<210> 16  
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 <212> DNA  
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<220>  
 <223> PCR primer HCG-ANTISENT

<400> 16  
 ttaagatttg tgataataac aagtactgca 30

<210> 17  
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 <223> glycalA - RT-PCR product

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 tatcccaactc cactaagggtc caagaagacg atgttggtcc aaaagaacgt cacctcagag 240  
 tccacttgct gtgttagctaa atcatataac agggtcacag taatgggggg tttcaaagtg 300  
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 agcatatccc actccactaa ggtccaagaa gacgatgttgc tccaaaaga acgtcacctc 240  
 agagtccact tgctgtgttag ctaaatcata taacagggtc acagtaatgg ggggtttcaa 300  
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 <213> Artificial Sequence

<220>  
 <223> PCR product AB-FSH  
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<210> 24  
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 <212> DNA  
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 <223> PCR product glycalwoTAA  
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 tccacttgct gtgttagctaa atcatataac agggtcacag taatgggggg tttcaaagtg 180  
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 gagaaccaca cggcgtgcca ctgcagtact tgttattatc acaaatct 300  
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<223> hybrid reverse primer ABLIGATION

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ttggtcagct cacagctatt agatttgtga taataacaag tactgcagtg g

51

<210> 26

<211> 368

<212> DNA

<213> Artificial Sequence

<220>

<223> PCR product glycalwoTAAUR

<400> 26

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60

gttctccatt cgcctcctga tgtgcaggat tgcccagaat gcacgctaca ggaaaaccca

120

ttcttctccc agccgggtgc cccaatactt cagtgcattgg gctgctgctt ctctagagca

180

tatcccactc cactaaggc caagaagacg atgttggtcc aaaagaacgt cacctcagag

240

tccacttgct gtgtagctaa atcatataac agggtcacag taatgggggg tttcaaagtg

300

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ctgaccaa

368

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<212> PRT

<213> Artificial Sequence

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<223> synthetic peptide AB-FSH

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35 40 45

Ile Leu Gln Cys Met Gly Cys Cys Phe Ser Arg Ala Tyr Pro Thr Pro  
50 55 60

Leu Arg Ser Lys Lys Thr Met Leu Val Gln Lys Asn Val Thr Ser Glu  
65 70 75 80

Ser Thr Cys Cys Val Ala Lys Ser Tyr Asn Arg Val Thr Val Met Gly  
85 90 95

Gly Phe Lys Val Glu Asn His Thr Ala Cys His Cys Ser Thr Cys Tyr  
100 105 110

Tyr His Lys Ser Asn Ser Cys Glu Leu Thr Asn Ile Thr Ile Ala Ile  
115 120 125

Glu Lys Glu Glu Cys Arg Phe Cys Ile Ser Ile Asn Thr Thr Trp Cys  
130 135 140

Ala Gly Tyr Cys Tyr Thr Arg Asp Leu Val Tyr Lys Asp Pro Ala Arg  
145 150 155 160

Pro Lys Ile Gln Lys Thr Cys Thr Phe Lys Glu Leu Val Tyr Glu Thr  
 165 170 175  
 Val Arg Val Pro Gly Cys Ala His His Ala Asp Ser Leu Tyr Thr Tyr  
 180 185 190  
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 Met Lys Glu  
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 <223> CDNA sequence of INF-beta without stop codon  
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 cactgtgcct ggaccatagt cagagtggaa atcctaagga acttttactt cattaacaga 240  
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 <223> INF-alpha-2B sequence with enterokinase site

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 tttggatttc cccaggagga gtttggcaac cagttccaaa aggctgaaac catccctgtc  
 ctccatgaga tgatccagca gatcttcaat ctcttcagca caaaggactc atctgctgtc  
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 180  
 240  
 300

gaaggcctgtg tgatacaggg ggtgggggtg acagagactc ccctgatgaa ggaggactcc 360  
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 agcccttgtg cctgggaggt tgtcagagca gaaatcatga gatcttttc tttgtcaaca 480  
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 <223> INF-beta/INF-alpha-2B sequence with enterokinase site

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 tatgagatgc tccagaacat ctttgctatt ttcagacaag attcatctag cactggctgg 300  
 aatgagacta ttgttgagaa ctcctggct aatgtctatc atcagataaa ccatctgaag 360  
 acagtccctgg aagaaaaact ggagaaagaa gatttcacca gggaaaact catgagcagt 420  
 ctgcacctga aaagatatta tggaggatt ctgcattacc tgaaggccaa ggagtacagt 480  
 cactgtgcct ggaccatagt cagagtggaa atcctaagga acttttactt cattaacaga 540  
 cttacagggtt acctccgaaa cgacgacgac gacaagtgtg atctgcctca aacccacagc 600  
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